



IFWO

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/764,604

DATE: 08/27/2004

TIME: 11:08:06

Input Set : A:\sequence.listing.ascii.txt  
 Output Set: N:\CRF4\08272004\J764604.raw

3 <110> APPLICANT: McLachlan, Karen  
 4 Glaser, Scott  
 5 Peach, Robert  
 6 Rowe, Anthony  
 8 <120> TITLE OF INVENTION: Compositions and Methods for Treating Cancer Using IGSF9 and  
 9 LIV-1  
 11 <130> FILE REFERENCE: 2159.0030001  
 13 <140> CURRENT APPLICATION NUMBER: US 10/764,604  
 14 <141> CURRENT FILING DATE: 2004-01-27  
 16 <150> PRIOR APPLICATION NUMBER: US 60/442,535  
 17 <151> PRIOR FILING DATE: 2003-01-27  
 19 <160> NUMBER OF SEQ ID NOS: 43  
 21 <170> SOFTWARE: PatentIn version 3.2  
 23 <210> SEQ ID NO: 1  
 24 <211> LENGTH: 3490  
 25 <212> TYPE: DNA  
 26 <213> ORGANISM: Homo sapiens  
 28 <400> SEQUENCE: 1  
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 31 cgagggaaagc ctgagggttgt atcgggtggt ggccgggctg aggagagtgt ggtgctggc 120  
 33 tggacactgc tgccccccggc cggccggccc cccctgcatt tcattcgatg gctgcgttt 180  
 35 ggattccctgc ttcccatctt catccagttc ggcctctact ctcccccata tgaccctgtat 240  
 37 tacgtggac gagtcggcgt gcagaagggg gcctctctcc agattgaggg tctccgggtg 300  
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 45 cgtggcagcc ccctgcctca tgtgacgtgg aagctccgag gaaaggaccc tggccagggc 540  
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 49 tctgggtct acacctgcac agcctccagc actgaggcca ggcacccca cggccacccag 660  
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 61 tgcattgcgg gggtgatccg ctggccgggtt cgtgccaacc cccactgtc ctttgtcagc 1020  
 63 tggaccaagg atggaaaggc cctgcagctg gacaagttcc ctggctggc ccaggcaca 1080  
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 67 ccctacaaca gtcattgtac cggccggccc tctcctgtga cccgcgtgtc gctcaaggct 1200  
 69 ccccccagctt ttatagagcg gccaaggaa gaatatttcc aagaagtagg gggggagctg 1260  
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 81 agagattcag tgtctggta accccactgg ccaagcgcc tgaccgaatg caccatgact 1620  
 83 gggtgtcctt ggcaagtgcct gtgggggttg ctacccctt agtgcacggg ctgcagcccc 1680  
 85 acacccaga ccagttcage gtgttagtc agaacaagct ggggagtggt cccttageg 1740  
 87 aaatcgcttt gtctgctccg gaagggttc ctaccaccc agtgcaccc gggcttcccc 1800  
 89 caacagagat accgcctccc ctgtccctc cgccgggtct ggtggcagtg aggacacccc 1860  
 91 gggggtaact cctgcattgg gatccccag agtggtccc taagagactg gatggctacg 1920  
 93 tcttggaaagg ccggcaaggc tccagggtt gggaggtgtt ggacccggct gtggcaggca 1980  
 95 cagaaaacaga gctgctggta ccaggcctca tcaaggatgt tctctacgag ttccgcctcg 2040  
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 99 gtctggaggt ctacccttcg cgcacgcgc tgccgggctt cctgcctcag cccgtgtgg 2160  
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 105 cacctcttat ttctctccg accgggaagt cagctgcacc ctctgtctg ggctcaggca 2340  
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 113 ccactgtggc ggccccccag gaaaggtagt gccgggagca ggcaaaacctt cggactccag 2580  
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 133 ctggggccagt ggccctgtaga gatggccccc aaggagcat gtgggtacag tcagcaagag 3180  
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 137 attgtctggag actttgcacc tgggcttggc cagctcccg ctcagacctg aagctgagac 3300  
 139 agagcttagt gtgaagactc cagaggaggg ctgccttctg aacactgccc atgttactgg 3360  
 141 ccctgaggcc cgctgtgtt cccttcggga ggaatttctg gcttccgc gccgcccggaga 3420  
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 149 <211> LENGTH: 1163  
 150 <212> TYPE: PRT  
 151 <213> ORGANISM: Homo sapiens  
 153 <400> SEQUENCE: 2  
 155 Met Val Trp Cys Leu Gly Leu Ala Val Leu Ser Leu Val Ile Ser Gln  
 156 1 5 10 15  
 159 Gly Ala Asp Gly Arg Gly Lys Pro Glu Val Val Ser Val Val Gly Arg  
 160 20 25 30  
 163 Ala Glu Glu Ser Val Val Leu Gly Cys Asp Leu Leu Pro Pro Ala Gly  
 164 35 40 45  
 167 Arg Pro Pro Leu His Val Ile Glu Trp Leu Arg Phe Gly Phe Leu Leu  
 168 50 55 60  
 171 Pro Ile Phe Ile Gln Phe Gly Leu Tyr Ser Pro Arg Ile Asp Pro Asp  
 172 65 70 75 80

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175 Tyr Val Gly Arg Val Arg Leu Gln Lys Gly Ala Ser Leu Gln Ile Glu  
 176 85 90 95  
 179 Gly Leu Arg Val Glu Asp Gln Gly Trp Tyr Glu Cys Arg Val Phe Phe  
 180 100 105 110  
 183 Leu Asp Gln His Ile Pro Glu Asp Asp Phe Ala Asn Gly Ser Trp Val  
 184 115 120 125  
 187 His Leu Thr Val Asn Ser Pro Pro Gln Phe Gln Glu Thr Pro Pro Ala  
 188 130 135 140  
 191 Val Leu Glu Val Gln Glu Leu Glu Pro Val Thr Leu Arg Cys Val Ala  
 192 145 150 155 160  
 195 Arg Gly Ser Pro Leu Pro His Val Thr Trp Lys Leu Arg Gly Lys Asp  
 196 165 170 175  
 199 Leu Gly Gln Gly Gln Gly Val Gln Val Gln Asn Gly Thr Leu Arg  
 200 180 185 190  
 203 Ile Arg Arg Val Glu Arg Gly Ser Ser Gly Val Tyr Thr Cys Gln Ala  
 204 195 200 205  
 207 Ser Ser Thr Glu Gly Ser Ala Thr His Ala Thr Gln Leu Leu Val Leu  
 208 210 215 220  
 211 Gly Pro Pro Val Ile Val Val Pro Pro Lys Asn Ser Thr Val Asn Ala  
 212 225 230 235 240  
 215 Ser Gln Asp Val Ser Leu Ala Cys His Ala Glu Ala Tyr Pro Ala Asn  
 216 245 250 255  
 219 Leu Thr Tyr Ser Trp Phe Gln Asp Asn Ile Asn Val Phe His Ile Ser  
 220 260 265 270  
 223 Arg Leu Gln Pro Arg Val Gln Ile Leu Val Asp Gly Ser Leu Arg Leu  
 224 275 280 285  
 227 Leu Ala Thr Gln Pro Asp Asp Ala Gly Cys Tyr Thr Cys Val Pro Ser  
 228 290 295 300  
 231 Asn Gly Leu Leu His Pro Pro Ser Ala Ser Ala Tyr Leu Thr Val Leu  
 232 305 310 315 320  
 235 Cys Met Pro Gly Val Ile Arg Cys Pro Val Arg Ala Asn Pro Pro Leu  
 236 325 330 335  
 239 Leu Phe Val Ser Trp Thr Lys Asp Gly Lys Ala Leu Gln Leu Asp Lys  
 240 340 345 350  
 243 Phe Pro Gly Trp Ser Gln Gly Thr Glu Gly Ser Leu Ile Ile Ala Leu  
 244 355 360 365  
 247 Gly Asn Glu Asp Ala Leu Gly Glu Tyr Ser Cys Thr Pro Tyr Asn Ser  
 248 370 375 380  
 251 Leu Gly Thr Ala Gly Pro Ser Pro Val Thr Arg Val Leu Leu Lys Ala  
 252 385 390 395 400  
 255 Pro Pro Ala Phe Ile Glu Arg Pro Lys Glu Glu Tyr Phe Gln Glu Val  
 256 405 410 415  
 259 Gly Arg Glu Leu Leu Ile Pro Cys Ser Ala Gln Gly Asp Pro Pro Pro  
 260 420 425 430  
 263 Val Val Ser Trp Thr Lys Val Gly Arg Gly Leu Gln Gly Gln Ala Gln  
 264 435 440 445  
 267 Val Asp Ser Asn Ser Ser Leu Ile Leu Arg Pro Leu Thr Lys Glu Ala  
 268 450 455 460  
 271 His Gly His Trp Glu Cys Ser Ala Ser Asn Ala Val Ala Arg Val Ala

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272	465	470	475	480
275	Thr Ser Thr Asn Val Tyr Val Leu Gly Thr Ser Pro His Val Val Thr			
276	485	490	495	
279	Asn Val Ser Val Val Ala Leu Pro Lys Gly Ala Asn Val Ser Trp Glu			
280	500	505	510	
283	Pro Gly Phe Asp Gly Gly Tyr Leu Gln Arg Phe Ser Val Trp Tyr Thr			
284	515	520	525	
287	Pro Leu Ala Lys Arg Pro Asp Arg Met His His Asp Trp Val Ser Leu			
288	530	535	540	
291	Ala Val Pro Val Gly Ala Ala His Leu Leu Val Pro Gly Leu Gln Pro			
292	545	550	555	560
295	His Thr Gln Tyr Gln Phe Ser Val Leu Ala Gln Asn Lys Leu Gly Ser			
296	565	570	575	
299	Gly Pro Phe Ser Glu Ile Val Leu Ser Ala Pro Glu Gly Leu Pro Thr			
300	580	585	590	
303	Thr Pro Ala Ala Pro Gly Leu Pro Pro Thr Glu Ile Pro Pro Pro Leu			
304	595	600	605	
307	Ser Pro Pro Arg Gly Leu Val Ala Val Arg Thr Pro Arg Gly Val Leu			
308	610	615	620	
311	Leu His Trp Asp Pro Pro Glu Leu Val Pro Lys Arg Leu Asp Gly Tyr			
312	625	630	635	640
315	Val Leu Glu Gly Arg Gln Gly Ser Gln Gly Trp Glu Val Leu Asp Pro			
316	645	650	655	
319	Ala Val Ala Gly Thr Glu Thr Glu Leu Leu Val Pro Gly Leu Ile Lys			
320	660	665	670	
323	Asp Val Leu Tyr Glu Phe Arg Leu Val Ala Phe Ala Gly Ser Phe Val			
324	675	680	685	
327	Ser Asp Pro Ser Asn Thr Ala Asn Val Ser Thr Ser Gly Leu Glu Val			
328	690	695	700	
331	Tyr Pro Ser Arg Thr Gln Leu Pro Gly Leu Leu Pro Gln Pro Val Leu			
332	705	710	715	720
335	Ala Gly Val Val Gly Gly Val Cys Phe Leu Gly Val Ala Val Leu Val			
336	725	730	735	
339	Ser Ile Leu Ala Gly Cys Leu Leu Asn Arg Arg Arg Ala Ala Arg Arg			
340	740	745	750	
343	Arg Arg Lys Arg Leu Arg Gln Asp Pro Pro Leu Ile Phe Ser Pro Thr			
344	755	760	765	
347	Gly Lys Ser Ala Ala Pro Ser Ala Leu Gly Ser Gly Ser Pro Asp Ser			
348	770	775	780	
351	Val Ala Lys Leu Lys Leu Gln Gly Ser Pro Val Pro Ser Leu Arg Gln			
352	785	790	795	800
355	Ser Leu Leu Trp Gly Asp Pro Ala Gly Thr Pro Ser Pro His Pro Asp			
356	805	810	815	
359	Pro Pro Ser Ser Arg Gly Pro Leu Pro Leu Glu Pro Ile Cys Arg Gly			
360	820	825	830	
363	Pro Asp Gly Arg Phe Val Met Gly Pro Thr Val Ala Ala Pro Gln Glu			
364	835	840	845	
367	Arg Ser Gly Arg Glu Gln Ala Glu Pro Arg Thr Pro Ala Gln Arg Leu			
368	850	855	860	

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371	Ala	Arg	Ser	Phe	Asp	Cys	Ser	Ser	Ser	Pro	Ser	Gly	Ala	Pro	Gln	
372	865			870			875								880	
375	Pro	Leu	Cys	Ile	Glu	Asp	Ile	Ser	Pro	Val	Ala	Pro	Pro	Pro	Ala	Ala
376				885				890							895	
379	Pro	Pro	Ser	Pro	Leu	Pro	Gly	Pro	Gly	Pro	Leu	Leu	Gln	Tyr	Leu	Ser
380				900				905							910	
383	Leu	Pro	Phe	Phe	Arg	Glu	Met	Asn	Val	Asp	Gly	Asp	Trp	Pro	Pro	Leu
384		915				920									925	
387	Glu	Glu	Pro	Ser	Pro	Ala	Ala	Pro	Pro	Asp	Tyr	Met	Asp	Thr	Arg	Arg
388		930				935								940		
391	Cys	Pro	Thr	Ser	Ser	Phe	Leu	Arg	Ser	Pro	Glu	Thr	Pro	Pro	Val	Ser
392		945				950									960	
395	Pro	Arg	Glu	Ser	Leu	Pro	Gly	Ala	Val	Val	Gly	Ala	Gly	Ala	Thr	Ala
396				965				970							975	
399	Glu	Pro	Pro	Tyr	Thr	Ala	Leu	Ala	Asp	Trp	Thr	Leu	Arg	Glu	Arg	Leu
400				980				985							990	
403	Leu	Pro	Gly	Leu	Leu	Pro	Ala	Ala	Pro	Arg	Gly	Ser	Leu	Thr	Ser	Gln
404		995				1000									1005	
407	Ser	Ser	Gly	Arg	Gly	Ser	Ala	Ser	Phe	Leu	Arg	Pro	Pro	Ser	Thr	
408		1010				1015									1020	
411	Ala	Pro	Ser	Ala	Gly	Gly	Ser	Tyr	Leu	Ser	Pro	Ala	Pro	Gly	Asp	
412		1025				1030									1035	
415	Thr	Ser	Ser	Trp	Ala	Ser	Gly	Pro	Glu	Arg	Trp	Pro	Arg	Arg	Glu	
416		1040				1045									1050	
419	His	Val	Val	Thr	Val	Ser	Lys	Arg	Arg	Asn	Thr	Ser	Val	Asp	Glu	
420		1055				1060									1065	
423	Asn	Tyr	Glu	Trp	Asp	Ser	Glu	Phe	Pro	Gly	Asp	Met	Glu	Leu	Leu	
424		1070				1075									1080	
427	Glu	Thr	Leu	His	Leu	Gly	Leu	Ala	Ser	Ser	Arg	Leu	Arg	Pro	Glu	
428		1085				1090									1095	
431	Ala	Glu	Thr	Glu	Leu	Gly	Val	Lys	Thr	Pro	Glu	Glu	Gly	Cys	Leu	
432		1100				1105									1110	
435	Leu	Asn	Thr	Ala	His	Val	Thr	Gly	Pro	Glu	Ala	Arg	Cys	Ala	Ala	
436		1115				1120									1125	
439	Leu	Arg	Glu	Glu	Phe	Leu	Ala	Phe	Arg	Arg	Arg	Arg	Asp	Ala	Thr	
440		1130				1135									1140	
443	Arg	Ala	Arg	Leu	Pro	Ala	Tyr	Arg	Gln	Pro	Val	Pro	His	Pro	Glu	
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461	tgtgacctgc	tgc	ccccccggc	cggccggccc	cccctgc	atgc	tcatcgatgc	gtcg	cgcttt						180	
463	ggattcctgc	ttcccatctt	catccagttc	ggcctctact	ctccccgaat	tgacc	ctgt								240	

**VERIFICATION SUMMARY**

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